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---MKKMSPYQNKNEYEILESSSNNTNTPNRYP---FANNRDMSTMS-----WNDCQGI 48
axmi008
              ------VEVL 24
crylAa
              -----VEVL 24
cry1Ac
              -----MKLKNQDKHQSFSSNAKVDKISTDS---LKNETDIELQNINHEDCLKMSEYE 49
crylIa
              -----MNNVLNSGRTTICDAYNVVAHDPFSFEHKSLDTIQKEWMEWKRTDHSLYVAPVV 54
cry2Aa
              MIRKGGRKMNPNNRSEHDTIKTTENNEVPTNHVQYPLAETPNPTLEDLNYKEFLRMTADN 60
cry3Aa1
              ------MNPNNRSEHDTIKVTPNSELQTNHNQYPLADNPNSTLEELNYKEFLRMTEDS 52
cry3Bb
              -----MNPYQNKNEYETLNASQKKLNISNNYTRYPIENSPKQLLQSTNYKDWLNMCQQN 54
cry4Aa
              -----MN-------SGYPLANDLQGSMKNTNYKDWLAMCENN 30
cry4Ba
              ------MIIDSKTTLPRHSLIHTIKLNSNKKYGPGDMTNGNQFIISKQEWATIGAYI 51
cry6Aa
              -----TQKALSPSLKNMNYQDFLSITERE 47
cry7Aa
              -----MSPNNQNEYEIIDATPSTSVSSDSNRYPFANEPTDALQNMNYKDYLKMSGGE 52
cry8Aa
              -----MNPYQNKNEYEIFNAPSNGFSKSNNYSRYPLANKPNQPLKNTNYKDWLNVCQDN 54
cry10Aa
              -----MHYYGNRNEYDILNASSNDSNMSNTYPRYPLANPQQDLMQNTNYKDWLNVCEGY 54
cry16Aa
              -----MNSYQNKNEYEILDAKRNTCHMSNCYPKYPLANDPQMYLRNTHYKDWINMCEEA 54
cry19Ba
              -----MNQYQNKNEYEILESSQNNMNMPNRYP---FADDPNAVMKNGNYKDWVNECEGS 51
cry24Aa
              -----MNPYQNKSECEILNAPLNNINMPNRYP---FANDPNAVMKNGNYKDWLNECDGI 51
crv25Aa
              -----NSYENKNEYEILNDSKK-SNMSNPYLRYPLANDSLASMQNTNYKDWLTMCDRT 52
cry39Aa1
               ------NSYENKNEYEILESSSNNTNMPNRYP---FANDRDMSTMS-----FNDCQGI 44
cry40Aa1
              S-----WDEIWESVETITSIGINLIEFVIEPSLGGINTLLSIIGKLIPTNRQT- 96
axmi008
              G-----GERIETGYTPIDISLSLTQFLLS-EF-VPGAGFVLGLVDIIWGIFGPSQ--- 72
cry1Aa
              G-----GERIETGYTPIDISLSLTQFLLS-EF-VPGAGFVLGLVDIIWGIFGPSQ--- 72
cry1Ac
              N-----VEPFVS-ASTIQTGIGIAGKILG-TLGVPFAGQVASLYSFILGELWP-KGK- 98
cry1Ia
              G------KRILSELWGIIFPSGSTN 88
cry2Aa
             N-----TEALDSSTTKDVIQKGISVVGDLLG-VVGFPFGGALVSFYTNFLNTIWPSE--- 111
cry3Aa1
             S----TEVLDNSTVKDAVGTGISVVGQILG-VVGVPFAGALTSFYQSFLNTIWPSDA-- 104
cry3Bb
              QQYGGDFETFIDSG---ELSAYTIVVGTVLTGFGFTTPLGLALIGFGTLIPVLFPAQDQS 111
cry4Aa
              QQYG-VNPAAINSS---SVSTALKVAGAILK--FVNPPAGTVLTVLSAVLPILWPTNTPT 84
cry4Ba
             Q-----TGLGLPVNEQQLRTHVNLSQDISIPSDFSQLYDVYCSDKTS- 93
cry6Aa
             O-----PEALASGNTAINTVVSVTGATLS-ALGVPGASFITNFYLKIAGLLWPENGK- 98
cry7Aa
             NPEL-FGNPETFIS-SSTIQTGIGIVGRILG-ALGVPFASQIASFYSFIVGQLWPSKSV- 108
cry8Aa
             QQYGNNAGNFASSETIVGVSAGIIVVGTMLG-AFAAPVLAAGIISFGTLLPIFWQGSDPA 113
cry10Aa
              -----HIENPREASVRAGLGKGLGIVSTIVGF-FGGSIILDTIGLFYQISELLWPEDDT- 107
cry16Aa
              ----SYASSGPSQL---FKVGGSIVAKILG-----MIPEVGPLLSWMVSLFWPTIEE- 99
cry19Ba
              N-----ISPSPAAAITSKIVSIVLKTLAKAVASSLADSIKSSLGISKTITENN- 99
cry24Aa
              T-----PSIFGTLGVLASIVISTIN-LATSPSIGDAFALVSSIGEYWPETKTS- 98
cry25Aa
               ----DTDVLSSRGA---VSTGVGMLSTILSL-FGIPLIGEGIDLLLGAADFLWPESDGG 103
cry39Aa1
               cry40Aa1
               ---VSALSICDLLSIIRKEVADSVLSDAIADF-DGKLKNYREYYLSYLGAWLKDGKPLQK 152
axmi008
               ----WDAFPVQIEQLINQRIEEFARNQAISRL--EGLSNLYQIYAESFREWEADPT---N 123
crylAa
              ----WDAFLVQIEQLINQRIEEFARNQAISRL--EGLSNLYQIYAESFREWEADPT---N 123
cry1Ac
              --NQWEIFMEHVEEIINQKISTYARNKALTDL--KGLGDALAVYHDSLESWVGNRN---N 151
cry1Ia
              ---LMQDILRETEQFLNQRLNTDTLARVNAEL--IGLQANIREFNQQVDNFLNPTQNPVP 143
cry2Aa
             --DPWKAFMEQVEALMDQKIADYAKNKALAEL--QGLQNNVEDYVSALSSWQKNPVSSRN 167
cry3Aa1
              --DPWKAFMAQVEVLIDKKIEEYAKSKALAEL--QGLQNNFEDYVNALNSWKKTPLSLRS 160
cry3Bb
              --NTWSDFITQTKNIIKKEIASTYISNANKIL--NRSFNVISTYHNHLKTWENNPNPQ-N 166
cry4Aa
             PERVWNDFMTNTGNLIDQTVTAYVRTDANAKM--TVVKDYLDQYTTKFNTWKREPNNQSY 142
cry4Ba
              -AEWWNKNLYPLIIKSANDIASYGFKVAGDPS--IKKDGYFKKLQDELDNIVDNNS---- 146
cry6Aa
               --I-WDEFMTEVEALIDQKIEEYVRNKAIAEL--DGLGSALDKYQKALADWLGKQD---D 150
cry7Aa
              --DIWGEIMERVEELVDQKIEKYVKDKALAEL--KGLGNALDVYQQSLEDWLENRN---D 161
cry8Aa
              N--VWQDLLN-IGGRPIQEIDKNIINVLTSIV--TPIKNQLDKYQEFFDKWEP-ARTHAN 167
cry10Aa
               QQYTWQDIMNHVEDLIDKRITEVIRGNAIRTL--ADLQGKVDDYNNWLKKWKDDPKSTGN 165
cry16Aa
               KNTVWEDMIKYVANLLKQELTNDTLNRATSNL--SGLNESLNIYNRALAAWKQNKNNFAS 157
cry19Ba
               ---VSQVSMVQVHQIINRRIQETILDLGESSL-NGLVAIYNRDYLGALEAWNNN-----K 150
crv24Aa
               ----FPLSVADVNRLIREALDQNAINRATGKFN-GLMDTYNTVYLKNLQDWYDTRIPANP 153
cry25Aa
               HQYTWEDLMNHIEELMDERLETEKRTTALDDL--RGLKALLGLFRDAFDSWEKNQNDPIA 161
cry39Aa1
               ---VSALSICDLLSIIRKEVADSVLSDAICRFLDGKLKNYREYYLPYLEAWLKDGKPLQK 149
cry40Aa1
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TNNSDI-GQLVYYFKLSERDFNEILGGSLSRN-----NAQVLLLPTFAQAANVQLLLLRD 206
axmi008
              PALR---EEMRIQFNDMNSALTTAIPLLAVQ-----NYQVPLLSVYVQAANLHLSVLRD 174
crylAa
              PALR---EEMRIQFNDMNSALTTAIPLFAVQ-----NYQVPLLSVYVQAANLHLSVLRD 174
cry1Ac
              TRAR---SVVKSQYIALELMFVQKLPSFAVS-----GEEVPLLPIYAQAANLHLLLLRD 202
crylIa
              LSIT-----SSVNTMQQLFLNRLPQFQIQG-----YQLLLLPLFAQAANMHLSFIRD 190
cry2Aa
              PHSQ---GRIRELFSQAESHFRNSMPSFAIS-----GYEVLFLTTYAQAANTHLFLLKD 218
cry3Aa1
              KRSQ---DRIRELFSQAESHFRNSMPSFAVS-----KFEVLFLPTYAQAANTHLLLLKD 211
cry3Bb
              TQDV---RTQIQLVHYHFQNVIPELVNSCPPNPSDCDYYNILVLSSYAQAANLHLTVLNQ 223
cry4Aa
              RTAV---ITQFNLTSAKLRETAVYFSNLVG------YELLLLPIYAQVANFNLLLIRD 191
cry4Ba
              -----DDDAIAKAIKDFKARCG-----ILIKEAKQYEEAAKNIVTSLD 184
cry6Aa
              PEAI---LSVATEFRIIDSLFEFSMPSFKVT-----GYEIPLLTVYAQAANLHLALLRD 201
cry7Aa
              ARTR---SVVSNQFIALDLNFVSSIPSFAVS-----GHEVLLLAVYAQAVNLHLLLLRD 212
cry8Aa
              AKAV---HDLFTTLEPIIDKDLDMLKNNAS------YRIPTLPAYAQIATWHLNLLKH 216
cry10Aa
              LSTL---VTKFTALDSDFNGAIRTVNNQGSP-----GYELLLLPVYAQIANLHLLLLRD 216
cry16Aa
              GELI---RSYINDLHILFTRDIQSDFSLG------GYETVLLPSYASAANLHLLLLRD 206
cry19Ba
              SNINYQ-TNVAEAFKTVEREFFTKLKGIYRTS----SSQITLLPTFTQAANLHLSMLRD 204
cry24Aa
              QGDSQLREAARRSLEEIERDFRKALAGEFAEAG-----SQIVLLPIYAQAANIHLLILKD 208
cry25Aa
              KNRV---GGYFEDVHTHFVKDMASIFSAT-----NYEVLLLPVYAQAANLHLLLLRE 210
cry39Aa1
              TNNSDI-GQLVKYFELSERDFNEILGGSLARN-----NAQILLLPYFCASCKCQLLLLRD 203
cry40Aa1
              AVQYKAQWFPFLSAENVRSELISPNSGCDFTGDYYERLKCKTAEYTNYCLYWYQVGLNQI 266
axmi008
              VSVFGQRWGFDAAT-----INSRYNDLTRLIGNYTDYAVRWYNTGLERV 218
cry1Aa
              VSVFGQRWGFDAAT------INSRYNDLTRLIGNYTDYAVRWYNTGLERV 218
cry1Ac
              ASIFGKEWGLSSSE-----ISTFYNRQVERAGDYSDHCVKWYSTGLNNL 246
crylIa
              VILNADEWGISAAT-----LRTYRDYLRNYTRDYSNYCINTYQTAFRGL 234
cry2Aa
              AOIYGEEWGYEKED------IAEFYKRQLKLTQEYTDHCVKWYNVGLDKL 262
cry3Aa1
              AQVFGEEWGYSSED-------VAEFYHRQLKLTQQYTDHCVNWYNVGLNGL 255
cry3Bb
              AVKFEAYLKNNROFDYLEP------LPTAIDYYPVLTKAIEDYTNYCVTTYKKGLNLI 275
cry4Aa
              G-----LINAQEWSLAR------SAGDQLYNTMVQYTKEYIAHSITWYNKGLDVL 235
cry4Ba
              QFLHGDQKKLEGVIN-----IQKRLKEVQTALNQAHGESSPAHKELLEKV 229
cry6Aa
              STLYGDKWGFTQNN-----IEENYNRQKKRISEYSDHCTKWYNSGLSRL 245
cry7Aa
              ASIFGEEWGFTPGE------ISRFYNRQVQLTAEYSDYCVKWYKIGLDKL 256
cry8Aa
              AATYYNIWLQNQGINPST-----FNSSNYYQGYLKRKIQEYTDYCIQTYNAGLTMI 267
cry10Aa
              AQIYGDKWWSARAN------ARDNYYQIQLEKTKEYTEYCINWYNKGLNDF 261
cry16Aa
cry19Ba
              VAIYG-KELGYPST-----DVEFYYNEQKYYTEKYSNYCVNTYKSGLESK 250
              AVMYQEGWNLQSHIN-------YSKELDDALEDYTNYCVEVYTKGLNAL 246
cry24Aa
              AMQFRTDLGLIRPVGVPITTS-----AEDPFESEFLLRIKKYTDHCISYYDDGLAKI 260
cry25Aa
              GVIYG-SRWGIAP-------AADFYHDQLLKYTAIYANHCVTWYNNGLAQQ 253
cry39Aa1
              AVQYEEQWFPFLSAENVRSELISPNSGCDFTGDYYERLKCKIAEYTDYCEYWYQAGLNQI 263
cry40Aa1
              KQGGTGAD----TWSKFNKFRREMTLAVLDIIAIFPTYDFEKYPLP-----THVELTR 315
axmi008
              WGPDSR-----TVSQLTR 265
crylAa
              WGPDSR-----DWVRYNQFRRELTLTVLDIVALFPNYDSRRYPIR----TVSQLTR 265
cry1Ac
              RGTNAE-----SWVRYNQFRRDMTLMVLDLVALFPSYDTQMYPIK-----TTAQLTR 293
crylIa
              NT-----RLHDMLEFRTYMFLNVFEYVSIWSLFKYQSLMVS-----SGA 273
cry2Aa
              RGSSYE-----SWVNFNRYRREMTLTVLDLIALFPLYDVRLYPKE-----VKTELTR 309
cry3Aa1
              RGSTYD-----AWVKFNRFRREMTLTVLDLIVLFPFYDIRLYSKG-----VKTELTR 302
cry3Bb
              KTTPDSNLDGNINWNTYNTYRTKMTTAVLDLVALFPNYDVGKYPIG-----VQSELTR 328
cry4Aa
              RNKSNG-----QWITFNDYKREMTIQVLDILALFASYDPRRYPADKIDNTKLSKTEFTR 289
cry4Ba
              KN-----FVVY 265
cry6Aa
              NGSTYE-----QWINYNRFRREMILMALDLVAVFPFHDPRRYSME-----TSTQLTR 292
cry7Aa
              KGTTSK-----SWLNYHQFRREMTLLVLDLVALFPNYDTHMYPIE-----TTAQLTR 303
cry8Aa
              RTNTNA-----TWNMYNTYRLEMTLTVLDLIAIFPNYDPEKYPIG-----VKSELIR 314
cry10Aa
              R-----TAGQWVNFNRYRREMTLTVLDIISMFPIYDARLYPT-----EVKTELTR 306
crv16Aa
              K-----QIG-WSDFNRYRREMTLSVLDIVALFPLYDTGLYPSK--DGKIHVKAELTR 299
cry19Ba
              R--GSTAI----DWLEFNSFRRDMTLMVLDLVAIFPNYNPVRYPLS-----TKISLSR 293
cry24Aa
              RSRGSDGE----TWWEFNKFRREMTLTVLDLVALYPTHNIKLYPIP-----TQTELSR 309
cry25Aa
              KE----LFAKSPNWNRFNAYRRDMTITVLDIIALFPTYDARLYTK-----PIKTELTR 302
cry39Aa1
              KOAGTGAD----TWAKFNKFRREMTLTVLDIIAIFQTYDFKKYPLP-----THVELTR 312
cry40Aa1
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EIYTDAVGYSS----GTYSWLRNWPNTFNGLEANGTRGP-GLVTWLSKIGIYNEYVSRY 369
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crylAa
               EIYTNPVLENFDGS-----FRG--SAQGIERSIRSP-HLMDILNSITIYTDAHRGY 313
cry1Ac
              EVYTDAIGTVHPHPSFTSTTWYNNNAPSFSAIEAAVVRNP-HLLDFLEQVTIYSLLS--R 350
cry1Ia
             NLYASGSGPQQTQS-----FTAQNWPFLYSLFQVNSNYILSGISGTRLSITFPNIG 324
cry2Aa
             DVLTDPIVGVNNLR------GYGTTFSNIEN-YIRKP-HLFDYLHRIQFHTRFQPGY 358
cry3Aa1
             DIFTDPIFSLNTLQ-----EYGPTFLSIEN-SIRKP-HLFDYLQGIEFHTRLQPGY 351
cry3Bb
             EIYQVLN---FEES-----PYKYYDFQYQEDSLTRRP-HLFTWLDSLNFYEKAQTTP 376
cry4Aa
             EIYTAL----VES-----PSSK-SIAALEAALTRDV-HLFTWLKRVDFWTNTIYQD 334
cry4Ba
             EILENTAVQHIKNQ------IDEIKKQLDSAQHDLDRDVKIIGMLNSINTDI 311
cry6Aa
             EVYTDPVSLSISNP------DIGPSFSQMENTAIRTP-HLVDYLDELYIYTSKYKAF 342
cry7Aa
              DVYTDPIAFNIVTSTGFCNPWSTHSGILFYEVENNVIRPP-HLFDILSSVEINTSRGGIT 362
crv8Aa
              EVYTNVN------SDTFRTITELENGLTRNP-TLFTWINQGRFYTRNSRDI 358
cry10Aa
              EIYSDVINGEIYGL-----MTPYFSFEKAESLYTRAP-HLFTWLKGFRFVTNS-ISY 356
cry16Aa
              EIYSDVINDHVYGL------MVPYISFEHAESLYTRRP-HAFTWLKGFRFVTNS-INS 349
cry19Ba
              KIYTDPVGRTDSPS-FGDWTNTGRTLANFNDLEREVTDSP-SLVKWLGDMTIYTGAIDSY 351
cry24Aa
               VVYTDPVGCFGNRK-----SDIFSRLNFDYLENRLTRPR-EPFNYLNSVQLFASTVSNS 362
cry25Aa
               EIYSDVLNLDVYGV------QQT--DLNKNEAAFTRSP-HLVTRLRGFDFYTRTKYAY 351
cry39Aa1
               EIYTDPVGYSS-----GTYSWLKYWTGAFNTLEANGTRGP-GLVTWLRSIGIYNEYVSRY 366
cry40Aa1
               FAG-----WVGTRHYEDYTKGNGIFQR-------MSGTTSNDLRNIDFQ---NADV 410
axmi008
               N-------YWSGHQITASPVGFSGPEFAFPL-----FGNAGNAAPP-VLVSLTGLGIF 358
cry1Aa
               Y-----YWSGHQIMASPVGFSGPEFTFPL-----YGTMGNAAPQQRIVAQLGQGVY 359
cry1Ac
               WSNTQYMN-MWGGHKLEFRTIGGTLNISTQ------GSTNTSINPVTLPFTSRDVY 399
crylIa
               GLPG----STTTHSLNSARVNYSGGVSSGLIG-----ATNLNHNFNCSTVLPPLSTPF 373
cry2Aa
               YGNDSFN--YWSGNYVSTRPSIGSNDIITSPF-----YGNKSSEPVQ--NLEFNGEKVY 408
cry3Aa1
              FGKDSFN--YWSGNYVETRPSIGSSKTITSPF-----YGDKSTEPVQ--KLSFDGQKVY 401
cry3Bb
              NN-----FFTSHYNMFHYTL-DNISQKSSV-----FGNHNVTDKLKS-LGLATNIYI 421
cry4Aa
             LR-----FLSANKIGFSYTN-SSAMQESGI-----YGSSGFGSNLTHQIQLNSNVYK 380
cry4Ba
              DN-----LYSQGQEAIKVFQKLQGIWATIG------AQIENLRTTSL 347
cry6Aa
              SHEIQPDLFYWSAHKVSFKKSEQSN-LYTTGI-----YGKTSGYISS-GAYSFHGNDIY 394
cry7Aa
             LNNDAYIN-YWSGHTLKYRRTADSTVTYTA-----NYGRITSEKNSFALEDRDIF 411
cry8Aa
              LDPYDIF--SFTGNQMAFTHTN-DDRNIIWGA-----VHGNIISQDTSKVFPFYRNKPI 409
cry10Aa
               WT-----FLSGGQNKYSYTN-NSSINEGSF-----RGQDTDYGGTSSTINIPSNSYV 402
cry16Aa
               WT------FLSGGENRYFLTHGEGTIYNGPF------LGQDTEYGGTSSYIDISNNSSI 396
cry19Ba
              RPT----SPGDRIGVWYGNINAFYHTGRTDVVMFRQTGDTAYEDPSTFISNILYDDI 404
cry24Aa
               NNGE-----VLRGNLNKIMFEGGWTASRSGDG----VTTGTPFSTMDWSYGWGYPRKHYA 413
cry25Aa
               WR------YLAGHTNYFSFTG-NGTIYSSSF-----N-NWYDTDMTKSTINIPDYANI 396
cry39Aa1
               FSG-----MSGTTSNDLRDISFP---NSDI 407
cry40Aa1
               YKITSLAIMNLVGETTARPEYRVSKADFRRVGGPDLNYDAGNNGLSRMTIESTFPLVLHS 470
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               RTLSSP---LYRRIILGSGPNNQELFVLDGTEFSFASLTTNLPSTIYRQRGTVDSLDVIP 415
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              RTLSST---LYRRPFN-IGINNQQLSVLDGTEFAYG-TSSNLPSAVYRKSGTVDSLDEIP 414
cry1Ac
              RTESLAGLNLFLTQPVNGVPRVDFHWKFVTHPIASDNFYYPGYAGIG--TQLQDSENELP 457
cry1Ia
               VRSWLDSGTDREGVATSTNWQTESFQTTLSLRCGAFSARGNSNYFPDYFIRNISGVPLVI 433
cry2Aa
               RAVANTNLAVWPS-AVYSGVTKVEFSQYNDQTDEASTQTYDSKRNVG-AVS-WDSIDQLP 465
cry3Aa1
               RTIANTDVAAWPNGKVYLGVTKVDFSQYDDQKNETSTQTYDSKRNNG-HVSAQDSIDQLP 460
cry3Bb
               FLLNVISLDNKY-LNDYNNISKMDFFITNGTRLLEKELTAGSGQITYDVNKNIFGLPILK 480
cry4Aa
               TSITDTSSP-----SNRVTKMDFYKIDGT---LASYNSNITPTPEGLRTTFFGFSTNE 430
cry4Ba
               QEVQDSDDADEIQIELEDASDAWLVVAQEARDFTLNAYSTNSRQNLPINVISDSCNCSTT 407
cry6Aa
               RTLAAPSVVVYPYTQN-YGVEQVEFYGVKGHVHYRGDNKYDL-----TYDSIDQLP 444
cry7Aa
               EINSTVANLANYYQKAYGVPGSWFHMVKRGTSSTTAYLYSKTHTALQGCTQVYESSDEIP 471
cry8Aa
               DKVEIVRHR-----EYSDIIYEMIFFSNSSE---VFRYSSNSTIENNYKRTDSYMIPKQT 461
cry10Aa
               YNLWTENYEYIYPWGDPVNITKMNFSVTDNNSSKELIYGAHRTNKPVVRTDFDFLTNKEG 462
cry16Aa
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cry19Ba
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cry24Aa
               EITSRSQALPGLNNSIHVIVGIDSFRAIGPGGQGDHTFSLPGGDMYDCGKVQINPLEDYR 473
crv25Aa
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cry39Aa1
               FKIESKAIMNLVGEINARPEYRVSRAEFSESTAFIYLYDAGNSGLSSMTITSKLPGIKNP 467
cry40Aa1
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crylAa	PQDNSVPPRAGFSHRLSHVTMLSQAAGAVYTLRAPTFSWQHRSAEFNNIIP	466
cry1Ac	PQNNNVPPRQGFSHRLSHVSMFRSGFSNSSVSIIRAPMFSWIHRSAEFNNIIA	467
cry1Ia	PEATGQPNYESYSHRLSHIGLISASHVK-ALVYSWTHRSADRTNTIE	503
cry2Aa	RNEDLTRPLHYNQIRNIESPSGTPGGARAYLVSVHNRKNNIYAANENGTMIHLA	487
cry3Aa1	PETTDEPLEKGYSHQLNYVMCFLMQGSRGTIPVLTWTHKSVDFFNMID	513
cry3Bb	PETTDEPLEKAYSHQLNYAECFLMQDRRGTIPFFTWTHRSVDFFNTID	508
cry4Aa	RRENQGNPTLFPTYDNYSHILSFIKSLSIPATYKTQVYTFAWTHSSVDPKNTIY	534
cry4Ba	NTPNQPTVNDYTHILSYIKTDVIDYNSNRVSFAWTHKIVDPNNQIY	
cry6Aa	NMTSNQYSNPTTNMTSNQYMI	
cry7Aa	PDGEPIHEKYTHRLCHATAIFKSTPDYDNATIPIFSWTHRSAEYYNRIY	
cry8Aa	LDRTVPVAESYSHRLSHITSHSFSKNG-SAYYGSFPVFVWTHTSADLNNTIY	
cry10Aa	WKNEEYGHTLSYIKTDNYIFSVVRERRRVAFSWTHTSVDFQNTID	
cry16Aa	TELAKYNDYNHILSYMLINGETFGQKRHGYSFAFTHSSVDPNNTIA	
cry19Ba	NGPTTYNDYNHILSYMLINGETFGQKRHGYSFAFTHSSVDRYNTIV	
cry24Aa	EGNANDYSHLLCDVKILQEDSSNICEGRSSLLSHAWTHASLDRNNTIL	
cry25Aa	NSDHWISDMMTINQSVQLASNPTQTFAFSALSLGWHHSSAGNRNVYV	
cry39Aa1	EKPLTYENYSHILSYMTS-AQHFGDKKIGYTFAWMHESVDFDNRVD	
cry40Aa1	EPSYRDYSHRLSNAACVGAGNSRINVYGWTHTSMSKYNLIY	508
axmi008	ANQITQIPAVKSYYLQNYLANAYTYVIKGT-HTGGDLIRFLRTKSEYNAVYAGGGIRL	567
cry1Aa	SSQITQIPLTKSTNLGSGTSVVKGPGFTGGDILRRTSPGQISTLRVNITA	
crylAc	SDSITQIPAVKGNFLFNG-SVISGPGFTGGDLVRLNSSGNNIQNRGYIEV	516
cry1Ia	PNSITQIPLVKAFNLSSGAAVVRGPGFTGGDILRRTNTGTFGDIRVNINP	553
cry2Aa	PEDYTGFTISPIHATQVNNQTRTFISEKFGNQGDSLRFEQSNTTA	532
cry3Aa1	SKKITQLPLVKAYKLQSGASVVAGPRFTGGDIIQCTENGSAATIYV	
cry3Bb	AEKITQLPVVKAYALSSGASIIEGPGFTGGNLLFLKESSNSIAKFKV	
cry4Aa	THLTTQIPAVKANSLGTASKVVQGPGHTGGDLIDFKDHFKITCQ	
cry4Ba	TDAITQVPAVKSNFLNATAKVIKGPGHTGGDLVALTSNGTLSGRMEIQCK	
cry6Aa	SHEYTSLPNNFMLSR	_
cry7Aa	PNKITKIPAVKMYKLDDPSTVVKGPGFTGGDLVKRGSTGYIGDIKATVNS	543
cry8Aa	SDKITQIPAVKGDMLYLGGSVVQGPGFTGGDILKRTNPSILGTFAVTVNG	572
cry10Aa	LDNITQIHALKALKVSSDSKIVKGPGHTGGDLVILKDSMDFR	548
cry16Aa	ANKITQIPVVKASSINGSISIEKGPGFTGGDLVKMRADSGLTM	551
cry19Ba	PDKIVQIPAVKTNLVGANIIKGPGHTGGDLLKLEYERFLSL	543
cry24Aa	PDEITQIPAVTAYELRGNSSVVAGPGSTGGDLVKMSYHSVWS	553
cry25Aa	YDKITQIPATKTVREHPMIKGPGFTGGDLADLSSNSDILQYDLRSD	566
cry39Aa1	PDKITQIPAVKGDYLQYGYVKQGPGHTGGDLVSMIRTDRLGI	539
cry40Aa1	PDKITQIPAVKAFDISDTGPGQVIAGPGHTGGNVVSLPYYSRLKIRL	555
	•	
		c 0.1
axmi008	IINNKTAGQSYRIRFRYAADKAAFFSVYLYPGGWGSNRFVSLEKSYSGNYDDLK	
crylAa	PLSQRYRVRIRYASTTNLQFHTSIDGRPINQGNFSATMSSGSNLQ	561
cry1Ac	PIHFPSTSTRYRVRYASVTPIHLNVNWGNSSIFSNTVPATATSLDNLQ	
crylla	PFAQRYRVRIRYASTTDLQFHTSINGKAINQGNFSATMNRGEDLD	598
cry2Aa	RYTLRGNGNSYNLYLRVSSIGNSTIRVTINGRVYTVSNVNTTTNNDGVNDNGAR	200
cry3Aa1	TP-DVSYSQKYRARIHYASTSQITFTLSLDGAPFNQYYFDKTINKGDTLT	608
cry3Bb	TLNSAALLQRYRVRIRYASTTNLRLFVQNSNNDFLVIYINKTMNKDDDLT	
cry4Aa	HSNFQQSYFIRIRYASNGSANTRAVINLSIPGVAELG-MALNPTFSGTDYTNLK	
cry4Ba	TSIFNDPTRSYGLRIRYAANSPIVLNVSYVLQGVSRGTTISTESTFSRPNNIIPTDLK	
cry6Aa	NSNLEYKCPENNFMIYWYNNSDWYNNSDWYNN	
cry7Aa	PLSQKYRVRVRYATNVSGQFNVYINDKITLQTKFQNTVETIGEGKDLT	57I
cry8Aa	SLSQRYRVRIRYASTTDFEFTLYLG-DTIEKNRFNKTMDNGASLT	610
cry10Aa	VRFLKNVSRQYQVRIRYATNAPKTTVFLTGIDTISVELPSTTSRQNPNATDLT	
cry16Aa	RFKAELLDKKYRVRIRYKCNYSSKLILRKWKGEGYIQQQIHNISPTYGAFSY	503
cry19Ba	RIK-LIASMTFRIRIRYASNISGQMMINIGYQNPTYFNIIPTTSRDYTELK	607
cry24Aa	FKVYCSELKNYRVRIRYASHGNCQFLMKRWPSTGVAPRQWARHNVQGTFSNSMR YDDRLTEDVPFRIRIRCASIGVSTISVDNWGSSSPQVTVASTAASLDTLK	616
cry25Aa	YDDRLTEDVPFRIRIRCASIGVSTISVDNWGSSSPQVTVASTAASLDILK NVY-FPQPLDYRIRIRYSTSSNGYLYIYSPNTKIVYLPPTTLVDGQPTFDPMD	591
cry39Aa1	IPASTNKNYLVRVRYTSTSNGRLLVERWSPSSIINSYFFLPSTGPGDSFG	605
cry40Aa1	TAWDIMVNIDAKAKIIDIDMCKMDACMDACDIIINDILLIDGIIGEGDQLG	000

axmi008	YSDFKFAEIITPPLPSSNIQMDVEMQANSFQSDVNVVLDKIEFLPSNTTTLEYEGER	
cry1Aa	SGSFRTVGFTTPFNFSNGSSVFTLSAHV-FNSGNEVYIDRIEFVPAEVTFEAEY	
cry1Ac	SSDFGYFESANAFTSSLGNIVGVRN-FSGTAGVIIDRFEFIPVTATLEAEY	616
crylla	YKTFRTVGFTTPFSFLDVQSTFTIGAWN-FSSGNEVYIDRIEFVPVEVTYEAEY	
cry2Aa	FSDINIGNIVASDNTNVTLDINVTLNSGTPFDLMNIMFVPTNLPPLY	633
cry3Aa1	YNSFNLASFSTPFELSGNNLQIGVTG-LSAGDKVYIDKIEFIPVN	652
cry3Bb	YQTFDLATTNSNMGFSGDKNELIIGAES-FVSNEKIYIDKIEFIPVQL	652
cry4Aa	YKDFQYLEFSNEVKFAPNQNISLVFNRSDVYTNTTVLIDKIEFLPITRSIREDREKQ	688
cry4Ba	YEEFRYKDPFDAIVPMRLSSNQLITIAIQPLNMTSNNQVIIDRIEIIPITQSVLDETENQ	644
cry6Aa		
cry7Aa	YGSFGYIEYSTTIQFPDEHPKITLHLSD-LSNNSSFYVDSIEFIPVDVNYAEKE	
cry8Aa	YETFKFASFITDFQFRETQDKILLSMGD-FSSGQEVYIDRIEFIPVDETYEAEQ	
cry10Aa	YADFGYVTFPRTVPNKTFEGEDTLLMTLYGTPNHSYN-IYIDKIEFIPITQSVLDYTEKQ	660
cry16Aa	LESFTITTTENIFDLTMEVTYPYGRQFVEDIPSLILDKIEFLPTN	648
cry19Ba	FEDFQLVDTSYIYSGGPSISSNTLWLDNFSNGPVIIDKIEFIPLGITLNQAQGYD	
cry24Aa	YEAFKYLDIFTITPEENNFAFTIDLESGGDLFIDKIEFIPVSGSAFEYEGKQ	
cry25Aa	YESFQYVSIPGNYYFDSAPRIRLLRQPGRLLVDRIEIIPVNFFPLS	662
cry39Aa1	FSAFRVVEVPASFRASVAGYTNFTIEAGFGPVYIDKIEFIPDNTTTLEYEGGR	644
cry40Aa1	YVDTLVTTFNQPGVEIIIQNLDTPINVDKVEFIPVNSTALEYEGKQ	651
axmi008	DLEKTKNAVNDLFTN	693
cry1Aa	DLERAQKAVNELFTSSNQIGLKTDVTDYHIDQVSNLVECLSDEFCLDEKQELSEKVKHAK	674
cry1Ac	NLERAQKAVNALFTSTNQLGLKTNVTDYHIDQVSNLVTYLSDEFCLDEKRELSEKVKHAK	
cry1Ia	DFEKAQEKVTALFTSTNPRGLKTDVKDYHIDQVSNLVESLSDEFYLDEKRELFEIVKYAK	711
cry2Aa		
cry3Aa1		
cry3Bb		
cry4Aa	KLETVQQIINTFYANPIKNTLQSELTDYDIDQAANLVECISEELYPKEKMLLLDEVKNAK	748
cry4Ba	NLESEREVVNALFTNDAKDALNIGTTDYDIDQAANLVECISEELYPKEKMLLLDEVKNAK	704
cry6Aa		
cry7Aa	KLEKAQKAVNTLFTE-GRNALQKDVTDYKVDQVSILVDCISGDLYPNEKRELQNLVKYAK	
cry8Aa	DLEAAKKAVNALFTNTKD-GLRPGVTDYEVNQAANLVECLSDDLYPNEKRLLFDAVREAK	728
cry10Aa	NIEKTQKIVNDLFVN	675
cry16Aa		
cry19Ba	TYDQNANGMYHQNYSNSGYNYNQEYNTYYQSYNN	682
cry24Aa	NIEKTQKAVNDLFIN	674
cry25Aa	EQENKSVDSLFIN	675
cry39Aa1	DLEKTKN-AVNDLFTN	659
cry40Aa1	SLEKAQDVVNDLFVK	666

axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	MSPMFTSSTKNTLKIETTDYEIDQAAISIECMSDEQNPQEKIMLWDEIKLAKQLSQSRNL VNFMLTSGAKNMLKLETTDYEIDQMANAIENMSGEQYSQEKMMQWHDIKYAKQLSQARNLMFISNIKNTLKIETTDYEIDQAAISIECMSNEHSSKEEMMLWDEVKQAKQLSWSRNLMFTNGTKNTLKIETTDYEIDQAAISIECMSDEHSPKEKMMLWDEVKRAKLLSQSRNL INTFYANPIKNTLQSELTDYDIDQAANLVECISEELYPKEKMLLLDEVKNAKQLSQSRNV VNALFTNDAKDALNIGTTDYDIDQAANLVECISEELYPKEKMLLLDEVKNAKQLSQSRNV : *: *: ***:*** * :* :* ** :: :: :: : : * ** :* :	60 57 57 60
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	LQNGDFSGNDWTFGNDIIIGSNNPIFKGKFLQMRGARDIYGTLFPTYICQKIDESKL LQNGDFEDLFSG-WTTSNQMSIQADNATFKGNYLHMSGARDIYGTIFPTYIYQKIDESKL LYNGDFEDVSNG-WKTSNTIEIRENSPVFKGHYLHMFGARDIDGTLFPTYIYQKIEESKL LQNGDFGDFYGNDWKFGNNIIIGSNNSIFKGNFLQMSGARDIYGTIFPTYIYQKIDESKL LQNGDFESATLG-WTTSDNITIQEDDPIFKGHYLHMSGARDIDGTIFPTYIFQKIDESKL LQNGDFESATLG-WTTSDNITIQEDDPIFKGHYLHMSGARDIDGTIFPTYIFQKIDESKL * ** ***::*: * ***** **::********	119 116 117 119
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	KPYTRYRVRGFVGSSKDLKLMVTRYGKEIDAIMNVPNDLAYMQPNPSCGDYRCESSS KPYTRYLVRGFVGSSKDLELMVMRYGKEIDTVMNVPNDIPYVPSMPVCNELYDGQQPYPN KPYTRYRVRGFVGSSKDLKLMVTRYGKEIDAMMNVPNDLAYMQPTPSCGDSRCESSS KPYTRYRVRGFVGSSKDLRLMVTRYGKEIDAMMNVPNDLAYMQPNPSCGDSRCESSS KPYTRYLVRGFVGSSKDVELVVSRYGEEIDAIMNVPADLNYLYPSTFDCEGSN KPYTRYLVRGFVGSSKDVELVVSRYGEEIDAIMNVPADLNYLYPSTFDCEGSN ****** ******************************	179 173 174 172
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	QYVSQGYPTP-TDGYAPDMYACPQNIDRKHVKCHDRHPFDFHIDTGEVDTNTNVGIDVLL RHVGYYNPMP-VSQPSYTSDTCQCTPGKKHVVCHDSHQFKFHIDTGEVDYNTNLGIWVLF RYVSQGYPTPVTDGYASGRYACQSNRGTKHVKCHDRHPFDFHIDTGELDTNTNVGIDVLF QYVSQGYPTP-TDGYAPDRYACPSSSDKKHVMCHDRHPFDFHIDTGELDTNTNVGIDVLF RCETSAVPAN-IGNTSDMLYSCQYDTGKKHVVCQDSHQFSFTIDTGALDTNENIGVWVMF RCETSAVPAN-IGNTSDMLYSCQYDTGKKHVVCQDSHQFSFTIDTGALDTNENIGVWVMF :	238 233 233 231
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	KISNPDGYATVGNLEVIEEGPLTGEALAHVKQKEKKWKQHMEKKRWETQQAYDPAKQAVD KISSPDGYATLDNLEVIEEGPVRGEAVTHVKQKEKKWNQQMEKKRMETKRVYDRAKQAVD KISNPDGYATLGNLEVIEEGPLTGEALTHVKQKEKKWKQHMEKKRWETQQAYDPAKQAVD KISNPDGYATLGNLEVIEEGPLTGEALTHVKQKEKKWKQHMEKKRWETQQAYDPAKQAVD KISSPDGYASLDNLEVIEEGPIDGEALSRVKHMEKKWNDQMEAKRSETQQAYDVAKQAID KISSPDGYASLDNLEVIEEGPIDGEALSRVKHMEKKWNDQMEAKRSETQQAYDVAKQAID ***.***::.****************************	298 293 293 291
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	ALFTNEQELHYHITLDHIQNADRLVQSIPYVYHNWLPNAPGMNYDVYQELNARIMQGY ALFTGEELNYDVTLSHIKNADDLVQSIPYVHNEWLPDFPGMNYDIYQELNARIMQAR ALFTNEQELHYHITLDHIQNADRLIQAIPYVYHAWLPDAPGMNYDGYQGLNARIMQAY TLFTNEQELHYHITLDYIQTLIDWYSRFPIYTMTGYRDAPGMNYDGYQGLNARIMQAY ALFTNVQDEALQFDTTLAQIQYAEYLVQSIPYVYNDWLSDVPGMNYDIYVELDARVAQAR ALFTNVQDEALQFDTTLAQIQYAEYLVQSIPYVYNDWLSDVPGMNYDIYVELDARVAQAR :***. : *::. ** : : *:. ** : : *:.**	355 351 351 351
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	NLYDARNVITNGDFTQGLQGWHATGNAAVQQMDGASVLVLSNWSAGVSQNLHAQDHHGYV YLYDARNVITNGDFAQGLQGWHAEGKVEVQQMNGTSVLVLSNWSAGVSQNLHVQHPHGYL NLYDARNVITNGDFTQGLTGWHAAGKAMVQQMDGASVLVLSNWSAGVSQNLHVQEHHGYM NLYDARNVITNGDFTKGLQGWHAAGKAAVQQIDGASVLVLSNWSAGVSQNLHAQDHHGYM YLYDTRNIIKNGDFTQGVMGWHVTGNADVQQIDGVSVLVLSNWSAGVSQNVHLQHNHGYV YLYDTRNIIKNGDFTQGVMGWHVTGNADVQQIDGVSVLVLSNWSAGVSQNVHLQHNHGYV ***:**:****:*****::*: ****. *:. ***::*.********	415 411 411 411

axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	LRVIAKKEGPGKGYVTMMDCNGKQETLKFTSCEEGYMTKTVEVFPESDRVRIEIGETEGT LRVSAKKEGSGKGYVTRMSCNGKQETLTFTSCDGGYMTKTVEVFPESDRVRIEIGETEGS LRVIAKKEGPGKGYVTMMDCNGNRETLKFTSCEEGYMTKTVEVFPESDRVRIEIGETEGT LRVIAKKEGPGKGYVTMMDCNGNQETLKFTSCEEGYMTKTVEVFPESDRVRIEIGETEGT LRVIAKKEGPGNGYVTLMDCEENQEKLTFTSCEEGYITKTVDVFPDTDRVRIEIGETEGS LRVIAKKEGPGNGYVTLMDCEENQEKLTFTSCEEGYITKTVDVFPDTDRVRIEIGETEGS *** ***** * . * : : * . * . * . * : * * . * * * *	475 471 471 471
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	FYIDSIELLCMQGYDNNNNLHTGNMYEQSYNGNYNQNTSDVYYQGYTNNYNQDSSNMYNQ FYIESIELICMNGYTSNNNQNMSNMYDQSYSGNYSQNTSDMYDQGGSVAKFEK FYVDSIELLCMQGYASNNNPHTGNMYGQSYNGNYNQNTSDVYHQGYTNNYNQNSSNMYNQ FYVDSIELLCMQGYASNNNPHTGNMYGQSYNGNYNQNTSDVYHQGYTNNYNQNSSNMYNQ FYIESIELICMNE	528 531 531 484
axmi008-orf2 cry19Aa-orf2 crybun2orf2 crybun3orf2 cry4Aa cry4Ba	NYTHNDDLHSGCTCNQGHNSGCTCNQGYNR 561 E529 NYTHNDDLHSGCTCNQGHNSGCTCSQG558 NYTHNDDLHSGCTCNQGHNSGCTCSQG558	